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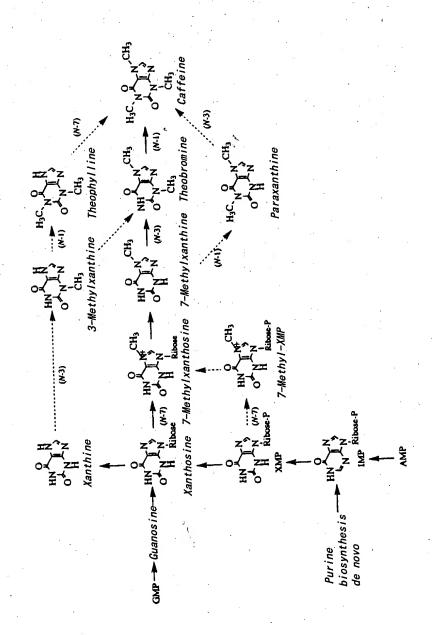
E: THEOBROMINE SYNTHASE POLYPEPTIDE

IT AND THE GENE ENCODING SAID POLYPEI

INVENTOR(S): HIROSHI SANO ET AL

APPLICATION SERIAL NO: UNASSIGNED

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F1G. 2

A GTCCTGCATA T ACTGGTTCTC G AGTTGCAGAT T GAAGAATGAA T CTCTATACCC A CCCCGAGGAG T CACTGCGAAC A CACATTITTA A GAATACCATG G AATCTATACA G GGCATCATTA A GAATGCAGCA A AAACCTTGT T TACTTTTTAT A AAAAAAAAAA	CCAAGGTGA TGGGATGCG TAGAACGTC WACTTGAGA CCCATGCATT WAAAGGAGCA GGATGCGTT WACTTACTTG WCTTCAGTAG CCTTCAGTAG CGCTTCTCTA ATTAGATCAG AGGTTACTG AGGTTACTG TTAGTTAGC TTAGTTAGC TTAGTTAGC TTAGTTAGC TTAGTTAGGT	AACCTGTCCT CTTCCGGACC CCACCATTCA AAGAAAATGG TITTACACTC TITTACACTC CGGAAGAGTT AGATGGCAAT AAGAAGTAAA TTGATGATGA TITTACGAACC GCTTGGGCAA TITTGTGTTA	TGAACAATGO AAACACACTT AGGOTTTTTCTC AGGCAAAATA TTCTTACAGT CAAAGCAAGT AAACGACTTC AGTGCAAGTA TTGCCAAGTA TTGCCAAGTA TTGCCTTCATT TGCGTTCTATT TGGGTTGTTT TGGGTTGTTT TGGGTTGTTT	THACCGTAT THACCGTT ACTGATCTT ACTGATCTT CCTCCGCCC GCCGAATG GTTGCTGAGG GTTGCTGAGG AGGAGGAGA AGATCCCATT AGTTCATTT TCTGATATCATTT TCTGATACGCC TCTGATATCATTT TCTGATACGC TTTGATACGC TTTGATACCC TTTGATACC TTTGATACCC TTTC	TIGTTGCGGGG GGGACACTG GGGACACTG TATCCCAGGG TATCCCAGGG TCCAGAAGG GTCCTTACTTG GGCGTCTGGG GTCCTTGGG GTCCTTGGG GTCCTTTGCG GGACAGCTTA TCCCCAGTATA GGGAAGGGT GGAAGGGTT GGAAGGGTT GGAAAGGGTT GGAAAGGGTT GGAAAGGGTT	C CAACTTGCCI T ACAAAGTAT TTCAATTCI TTCAATTCI TCCCAGCGGT TCCCAGCGGT TCCCAGCGGT TCCCAGCGGT TCCCAGCGGT TCCCAGCGGT TCCCAGCGGT TATATTGAA ACAGTGAAAAA ACAGTGATGAA TATACCTGAC TGCCAAAAAA CAGTGGGGT	C AACATCAAC G GACAAAGTT G GTTTTCATG C TCTTTCCAC T TTGGTGACT T CAATTTACG A GGAGATGAA TTGGACAGT TTGCAGACT TTGCAGACT A CATGCTAGAI A CCAGAGAGG GGGGTTCTA	G GCAGACTCTT G AATTGGGGAT A AAGATTTAC T GCGACGCCC T TCAATGTTCC T TTAAGCTCCG G CAGCGCATGT A GGTTTGCGAC T CAGACATATA T CCGAGTATTG	9 18 27 36 45 54 63 72 81 90 99 108 117 126 136
B TITAGCAGTC CO AAGCTACGCC AA CAACTTGCCC AA ACAAAGTATT GA	AGAATTCAT ACATCAACA ACAAAGTTA	CCTTCAATGA AGTGCATTAA GGCAAGAAAT	ACTGGTTCTC AGTTGCGGAT GAAGAATGAA	GCCAAGGTGA TTGGGATGCG TTAGAACGTC	AACCTGTCCT CTTCCGGACC CCACCATTCA	TGAACAATGC AAACACACTT GGTTTTTCTG	GTAGGGGAAT TTAACAGTTC ACTGATCTTT	TGTTGCGGGC GGGACATTGT TCCAAAATGA	98 186 276 360
TITCAATICG GI AATGCCTGGC TC TCCCAGCGGT TI ATAITTIGGAT CA CATTITGCAAA GC GGAAGAAAAA TI GGAAGAAAAA TI CAGCGATGAA CA TATACCTGAC AT	TTTTCATGT CTTTCCACG TGGTGACTG ATTTACGA GAGATGAAT TGGACAGTT TGGACAGTT	TGCTGCCAAG GCAGACTCTT AATTGGGGAT AAGATTTTAC TCGACGGCCC TCAATGTTCC TTAAGCTCCG	TTTCTACCGC CCCCGAGGAG CACTGCGAAC CACATTTTTA GAATACCATG AATCTATGCA TTATGATGCT	AAACTTGAGA TCAATGCATT AAAAGGAGCA AGGATTCGTT GACTTACTTG GCTTCAGTAG GGCTTCTCTA	AAGAAATGG TTTTACACTC TTTACTCTTC CGGAAGAGTT AGATGGCAAT AAGAATTAAA TTGATGATGA	ACGCAAGATA TTCTTACAGT CAAAGCAAGT GCTTTCACGC AAACGACTTG GTGCATAGTT TTGCCAAGTA	GGATCGTGCC CTTCAATTIT CCTCCGCCCG GGCCGAATGC GTTGTTGAGG GAGGAGGAAG AGATCCCATT	TAATAGCCGC TATCCCAGGT TCCAGAAGGC TCCTTACTTG GACATCTGGA GTTCTTTTGA CCCCAGAATA	450 540 630 720 810 900 990
TGCCAAAAAA CC TAGTGCGGTT GG	CAGAGAAGT	CAGACATATA	AAAGCTTGTT	TATAGTTGGT	TTTGTGCTA	TGGTTGTT	TCTGATACGG	GGAAAGGATT	1170 1260 1304
CTHTGGCAGT CC CAAGCTACGC CA CCAACTTGCC CA CCAACTTGCC CA TCCAAAGTAT TG ATTCCAATC GG CAATGCCGG TT CCTAGCGG TT CCATTGTAA AG AGGAAGAAAA AT AAATTTTATA CC CCGTTAGAGC AG CAAAGGTTCT CC TTTTTGTGTT GG TGGGTGTTAA AG TGGTGTTAAA	AGAATTCA AACAAGTT AACAAGTT TITTTCAAG TITTTCAAG TITTTCAC AATTTACG GAGTTGAA TITGGATAGT TITGGAGATTGAA TITGGATAGT TITGAGAGACT AATTTACGAA TITGGAGACT AATTTACGAA TITGGAGACT TITG	GCCTACAATC AAGTGCATTA GGCCAGGAAA ATTGCTGCCAA AGCAGACTCT GAATTGGGGA AAAGATTTTA ATTAGACGCC TTCAATCTTC TTTAAGGTCC AAAGGCTTCT AAAGGCTTCT AAAGGCTCCT AAAGGCCC AAAGGCTTCT AAAGGCCC AAAGGCCTCTTCT AAAGGCCC AAAGGCCTTCTT AAAGGCCC	AACTGGTTCT AAGTTGCGGA AGGAAGAATGA GCTTCTACCG YCCCCGAGGA TCAGTACGAA CCACATTTCT GGAATGCCAT CAGTCTATAT TTTACGATGC CAAGTCATTA TTTACGATGC TAGGATGCTATAT TTTACGATGC TAGGATGCTATAT TTTACGATGC TAGGATGCTATAT TTTACGATGC TTGGGGGTCT TTGGGGGGTCT	CGCCAAGGTG TTTGGGATGC ATTAGAACGT CAAACTTGAG GTCCATGCAT CAAAGGAGC AAAGGATTCAT ACCTTCAGCA TGGCTTCTCT TATCAGGAAGCT TATCATTCT TTCGGGTATT TTCGGGTATT TTCGGGTATT	AAACCTGTCC GCTTCTGGAC CCCACCATTC AAAGAAAATG TITTTACTCTT TCGGAAGAGT GAGATGGCAA GAAGAAGTAA ATTGACGATG ATTATACCTG CTCGCCAAAA GTGCTTTTTA	TTGAACAATG CAAACACACT AGATTTTCT GACGCAAAAT CTTGTTACTG CCAAAGCAAG TGTTTTCACA TAAACGACTT AGTGCATAGT AGTATTAAA ACATATTCAA ACCAGAGAA	CGTACGGAA TTTAACAGTT GAATGATCTTG GATGGTCGTGC TCTTCAATGG TCGTCTGCCC TGGCCGAATG GGTTGTTGAG TCGGAGGAA AGCAGAGTAT CAGGTTTGCG GTCAGACGTG	TTGTTGCGGG CGGGACATTG TTCCCAAATG CTAATAGGGG TTATCTCAGG GTCCAGAAGG CTCCTTACTT GGACATCTGG GGTTCTTTTG GTTGCATCTT AAGCATCTGT TAAAAAGTTTG	98 189 270 369 459 549 639 720 819 990 1088 1170 1260 1316
D				,					
AGCAGTICGA AT AGAIGAGA CA CAAGTIGGCAGA CAAGTIGGCAGA CAAGTIGGATA CAAGTIGGATA CAAGTIGAATTA GAATTICAAC AGAITTAAAT AGACACAAGA GTAGTATTAAT AGATTITAAT AGATTITAAT AGATTITAAT AGATTITAAT AGATTITAATA ATTITAATACA AATTIAATACA AATTIAATATAAATAAATAAATAATAAATAAATAAATA	ACAATCTGG TTAAAGTTG AAGAGAAGA ACAGCTTCT TCTTCCCCG GGATTGGTG TTACCACAT TACCGAATC TCCCATTATG GGATCACTTT TCTCCCACA TTTCTCCACAC TTTCTCCACAC TTTCTCCACAC TTTCTCCACAC TTTCTCCACAC	CTCTTGCCAA CGGATTTGGG ATGAATTAGA ACCGCAAACT AGGAGTCCAT, CAAACAAAGG TTCTAAGGAT CCCTAGACTT TTACACCTTC ATGCTGCCTT ACGAACCAAGG TGGCAAAGG TGTGCCGTTG	GGTGAAACCT ATGCGCTTCT ACGTCCCACC GCATTITTTG GAGTATTTAC TCATTCGAAA ACTTGACATG AGCAGAAGAA CCTCAGCAAGT CCTCGCAAGT CCTCGCAAGT CTGCTATAAT GGGGTCTTTC	TTCCTTGAAC GGACCAAACA ATTCAGATTT AATGGACGCA CACTCTTGTT TCTTCCAAAG GAGTTGTTTT GCAATAAACG GTAAAGTGCA GTAAAGTGCA CATTTTGGAG AATCTTACA	AATGCATACG CACTITTAAC TTCTGAATGA AGATAGGATC ACAGTGTTCA GATGTCGTCC CACGTGGCCG ACTTGATTGT TAGTTGAGGA CAGTAAGATC AAGCTATTAT TTTCTCTCGC	AGAATTGTTG AGTGCGGGAC TCTTTTCCAA GTGCCTAATA TTGGTTATCT GCCCGTCCAG AATGCTCCTT TGAGGGACTT GGAAGGTTCT CCATGAACAA GCCTGAACTA CAAAAAGCCA	CGGGCCAACT ATTGTGCAAA AATGATTTCA AGGGCAATGC CAGGTTCCCA AAGGCATATT ACCTGCATTT CTGGAGGAAG TGCGAAATTT ATTAAAGCAG TTCCACAGGC GAGAAGTCAG	TGCCCAACAT GTATTGACAA ATTCGGTTTT CTGGCTCTTT GCGGTTTGGT TGGATCAATT GTAAAGTAGA AAAAATTGGA TATATCTGGA AGTATGTGGC TTGCGAAGCA ACGTGTAAAA	98 188 270 360 450 540 630 720 810 990 1080 1170 1260 1298

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FIG. 3

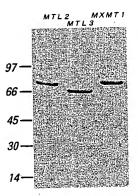
MXMT1	MELOEVLANDEGEGDTSYAKUASYN-LALAKVKPFLEQCTRELLRANLEN	49
MTL1	::::::::::::::::::::::::::::::::::::::	50
MTL2	::::::::::::::::::::::::::::::::::::::	50
MTL3	:::::::R::G:::::::::::SA::Q:V::::::V::::V:::::	50
MXMT1	INKCIKVADLOCASCENILLIVEDIVQSIDKVOQEEKNELEREPIQIFIN	99
MTL1	::::::::::::::::::::::::::::::::::::::	100
MTL2	::::::::::::::::::::::::::::::::::::::	100
MTL3	::::::::::::::::::::::::::::::::::::::	100
MXMT1	DLFONDENSVFKILLPSFYRKILÞKÐNERKTGSCLLSAMFGSFYGRILFPÐES	149
MTL1	- ::::::::::::::::::::::::::::::::::::	150
MTL2	::::::::::::::::::::::::::::::::::::::	150
MTL3	:::P::::::::::::::::::::::::::::::::::	150
MXMT1	MHFLHSCYSVHALSQVPSGLVIELGIGANKSSIYSSKGCRPPVQKAYIDQ	199
MTL1	:::::S::LQF::::::T::::T::::R::::::ASP::::::::	200
MTL2	::::::S::LQF::::::T::::R:::::ASP:::::::	200
MTL3	::::::::CLQ::::::::T:::::ST:::::::::::AS:L::::::::	200
MXMT1	FIKDFTIFIRIHSKELFSROMLICICKVÖEFDERNFIDLLDMAINILI	
MTL1	::::::::::::::::::::::::::::::::::::::	
MTL2	:::::::::::::::R:E:::L:::::::::::::::::	
MTL3	::::::::::::::::::::::::::::::::::::::	250
MXMT1	VEGLIEBEKLISHNIPFFIPSADEVKCIVEBEGSCEILYLEIFKAHYDAA	
MTL1	A::R:G::::::V:IY:A:V:::::M::::::F:::::Q:::IR:::G	
MTL2	:::H::::::::::::::::::::::::::::::::::	
MTL3	:::H:::::::L:VVI::::::::F::::::VL:::G	300
MXMT1	FSIDDDYFVRSH	
MTL1	::::::Q::::SFVYSD:HAR:AH::::::::::::::::::::::::::::::::	
MTL2	::::::Q::::SPEYSD:HAR;AH::::L::::::::N:::::I::I	
MTL3	::::::::::::::::::::::::::::::::::::::	337
MXMT1	FHRIAKPAKVIHMEKOCYNNLIISIAKKPEKSDV 378	٠.
MTL1	:::F:TN::::IRL:::F::::::::::1 365	
MTL2	:::F:TN:::::TRL:::F:::::::::::I 385	
MTL3	:::F::::::::::::::::::::::::::::::::::	

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FIG. 4



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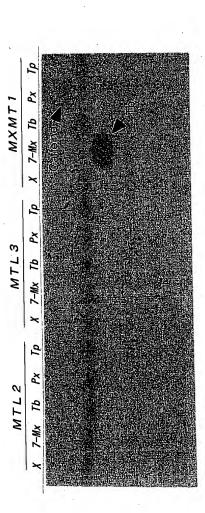
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INVENTOR(S): HIROSHI SANO, ET AL

APPLICATION SERIAL NO: UNASSIGNED

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FIG. 6

